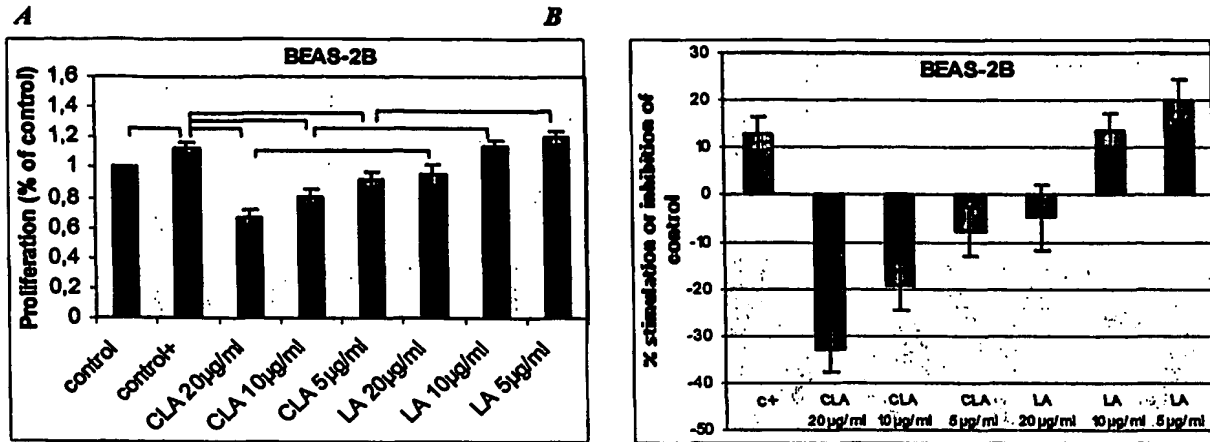
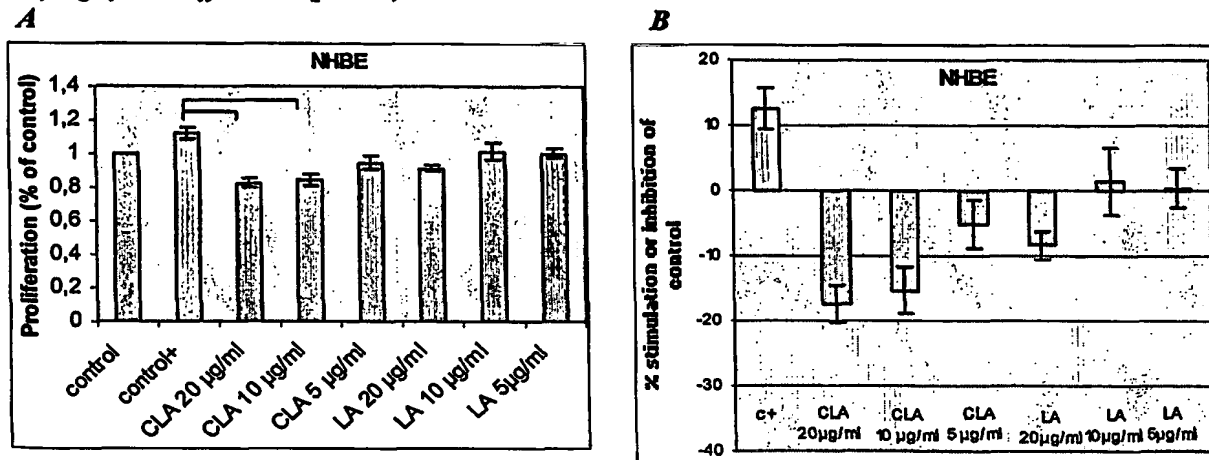


Figures

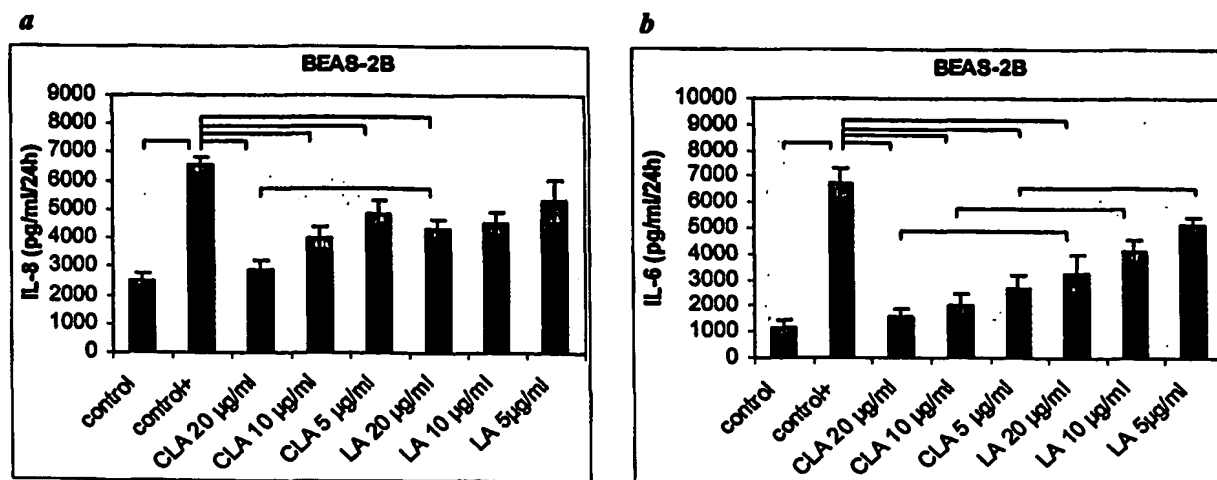
Figures 1A and 1B



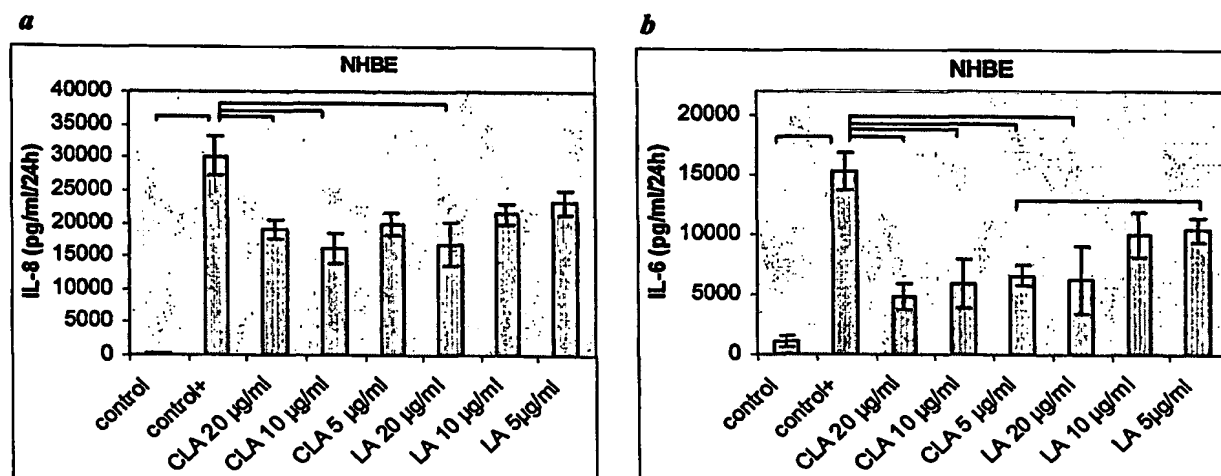
10 **Fig. 1A:** Modulation of LPS/serum-stimulated BEAS-2B by cis-9,trans-11 CLA and LA. The cells were incubated with the fatty acids at increasing concentrations for 24 h. (A) shows relative cell numbers compared with the unstimulated control (= 1) (B) depicts relative stimulation or inhibition observed. Data are means \pm SEM of 6 independent experiments performed at different days ($n = 6$). Connection of bars represents data with statistically significant differences ($p < 0.05$)



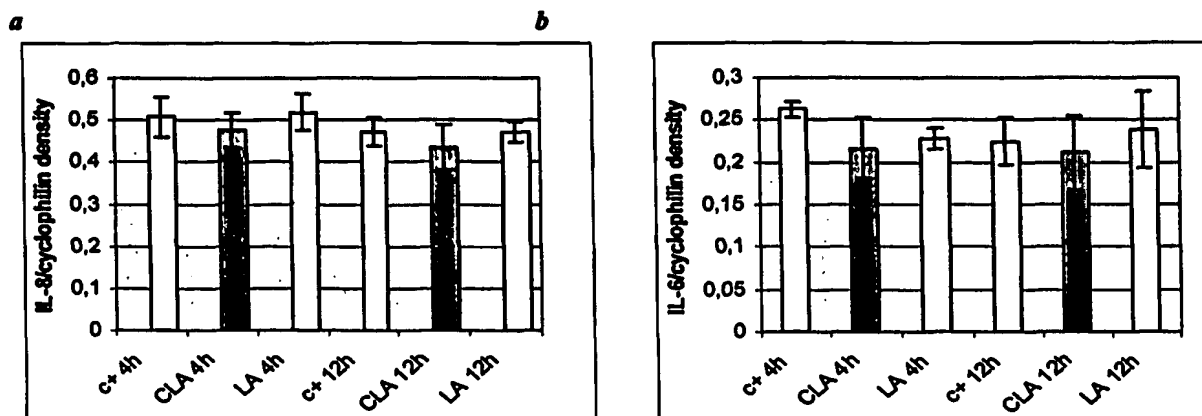
15 **Fig. 1B:** Modulation of LPS/serum-stimulated NHBE by cis-9,trans-11 CLA and LA. The cells were incubated with the fatty acids at increasing concentrations for 24 h. (A) shows relative cell numbers compared with the unstimulated control (= 1) (B) depicts relative stimulation or inhibition observed. Data are means \pm SEM of 6 independent experiments performed at different days ($n = 6$). Connection of bars represents data with statistically significant differences ($p < 0.05$)

Figures 2A and 2B

5 **Fig. 2A:** Effects of *cis*-9,*trans*-11-CLA and LA on the production of IL-8 (a) and IL-6 (b) by stimulated BEAS-2B after 24 h. Cells were incubated in the presence of 5 µg LPS and 10 % serum and different concentrations of either *cis*-9,*trans*-11-CLA or LA for 24 h and the supernatants were assessed for the concentration of cytokines. Data are shown as means ± SEM (n = 6), connection of bars represents data with statistically significant differences ($p < 0.05$).



15 **Fig. 2B:** Effects of *cis*-9,*trans*-11-CLA and LA on the production of IL-8 (a) and IL-6 (b) by stimulated BEAS-2B after 24 h. Cells were incubated in the presence of 5 µg LPS and 10 % serum and different concentrations of either *cis*-9,*trans*-11-CLA or LA for 24 h and the supernatants were assessed for the concentration of cytokines. Data are shown as means ± SEM (n = 6), connection of bars represents data with statistically significant differences ($p < 0.05$).

Figure 3

- 5 **Fig. 3: The expression of IL-8 (a) and IL-6 mRNA (b) in LPS- and serum-stimulated BEAS-2B over a 4-h and 12-h period of treatment without (c+) or with 20 μ g cis-9,trans-11-CLA or LA/mL. Total RNA was extracted and processed for RT-PCR. Cyclophilin was used as housekeeping gene. Data are shown as means \pm SEM (n = 4).**